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Research Article

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α -Ketoglutarate regulates acid-base balance through an intrarenal paracrine mechanism

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Paracrine communication between different parts of the renal tubule is increasingly recognized as an important determinant of renal function. Previous studies have shown that changes in dietary acid-base load can reverse the direction of apical α -ketoglutarate (α KG) transport in the proximal tubule and Henle's loop from reabsorption (acid load) to secretion (base load). Here we show that the resulting changes in the luminal concentrations of α KG are sensed by the α KG receptor OXGR1 expressed in the type B and non-A–non-B intercalated cells of the connecting tubule (CNT) and the cortical collecting duct (CCD). The addition of 1 mM α KG to the tubular lumen strongly stimulated Cl⁻-dependent HCO₃⁻ secretion and electroneutral transepithelial NaCl reabsorption in microperfused CCDs of wild-type mice but not *Oxgr1*^{-/-} mice. Analysis of alkali-loaded mice revealed a significantly reduced ability of *Oxgr1*^{-/-} mice to maintain acid-base balance. Collectively, these results demonstrate that OXGR1 is involved in the adaptive regulation of HCO₃⁻ secretion and NaCl reabsorption in the CNT/CCD under acid-base stress and establish α KG as a paracrine mediator involved in the functional coordination of the proximal and the distal parts of the renal tubule.

Introduction

α -Ketoglutarate (α KG) is an intermediate of the citric acid (TCA) cycle, an important anaplerotic substrate and a cofactor in a variety of enzymatic reactions. In addition to its direct effects on metabolic pathways, α KG was identified as a natural ligand to a GPCR, namely GPR99, which is also known as 2-oxoglutarate receptor 1 (OXGR1) (1). OXGR1 belongs to a cluster of so-called “metabolic” GPCRs, which also includes receptors for succinate (GPR91), lactate (GPR81), 3-hydroxy-octanoate (GPR109B), nucleotides (P2Y), fatty acids (FFAR), lipids (P2RY, CysLT, Oxer1, etc.), phospholipids (PAF), protease-activated receptors (PAR), and several orphan receptors (2). He et al. have shown that OXGR1 is a Gq-coupled GPCR that is predominantly expressed in distal tubules in the kidney (1). However, the functional role of OXGR1 has not been studied.

Previous studies in rats demonstrated that renal handling of α KG changes significantly in response to changes in acid-base status (3–5). α KG is freely filtered in the glomerulus and, under normal conditions, actively reabsorbed in the proximal tubule and Henle's loop. Acid load further stimulates α KG reabsorption, thus resulting in a drop in urinary output of α KG. Under base loading conditions, the blood concentration of α KG rises and net α KG reabsorption in the proximal tubule and Henle's loop is converted to net α KG secretion in the same nephron segments (3–5). This results in a significant increase in the urinary excretion of α KG. It has been proposed that excretion of α KG and other organic anions (e.g., citrate) in the urine represents the loss of “potential HCO₃⁻,” which provides the advantage of minimizing bicar-

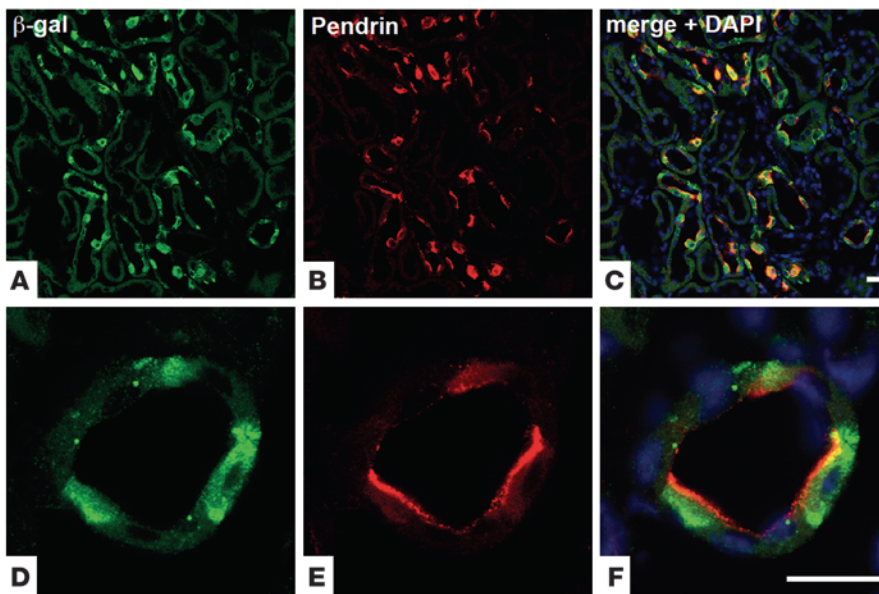
bonaturia under alkali load (6). The latter is important because it allows the excretion of base at a lower urinary pH, thereby diminishing the risk of nephrolithiasis due to the formation of calcium-phosphate precipitates [α KG: pKa₁(1.9), pKa₂(4.4); bicarbonate: pKa₁(6.1); HPO₄²⁻: pKa₂(6.7–6.8)] in the urine (5, 7). Collectively, these results demonstrated that acid-base status is a major factor determining blood levels of α KG and the rate of α KG excretion into urine. Importantly, Ferrier et al. have shown that there is no net transport of α KG beyond the beginning of the distal tubule accessible to micropuncture (3). This indicated that variations in the urinary α KG concentration are directly proportional to the variations in the luminal levels of α KG in the connecting tubule/cortical collecting duct (CNT/CCD), in which OXGR1 is expressed (see below). Taken together, these data led us to hypothesize that OXGR1 could be involved in the apical and/or basolateral sensing of acid-base status through the sensing of α KG concentrations in the tubular fluid and/or in the blood.

Testing this hypothesis revealed that luminal OXGR1 regulates Cl⁻-dependent HCO₃⁻ secretion and electroneutral transepithelial NaCl reabsorption in the type B and non-A–non-B intercalated cells of the CNT/CCD. We show that this regulation is functionally important since mice devoid of OXGR1 exhibited a reduced capacity to maintain acid-base equilibrium under base load conditions. We hypothesize that OXGR1-mediated NaCl reabsorption in the type B and non-A–non-B intercalated cells is required to compensate for the increased or decreased activity of sodium-hydrogen exchanger 3 (NHE3) in the proximal tubule and Henle's loop under acid or base loading conditions, respectively. Collectively, our results show for the first time that the kidney possesses a paracrine mechanism involved in the control of acid-base balance. In this mechanism, α KG acts as a paracrine mediator between the proximal and the distal parts of the renal tubule.

Authorship note: Natsuko Tokonami and Luciana Morla contributed equally to this work.

Conflict of interest: The authors have declared that no conflict of interest exists.

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**Figure 1**

OXGR1 is expressed in type B and non-A–non-B intercalated cells of the CNT and the CCD. (A) Immunostaining with anti- β -galactosidase antibody (green) shows that OXGR1 is expressed in a subset of cells in the renal cortex. (B) Immunolocalization of pendrin (red) in the apical membrane of type B and non-A–non-B intercalated cells. (C) Coimmunostaining with anti- β -galactosidase and anti-pendrin antibodies demonstrates that OXGR1 and pendrin are expressed in the same cells. Higher-magnification views of images in A–C are shown in D–F, respectively. Nuclei (DNA) are stained with DAPI (blue, C and F). Scale bars: 50 μ m.

Results

To identify renal cell types expressing OXGR1, we used mice whose *Oxgr1* gene was deleted and replaced with a reporter gene encoding β -galactosidase (*LacZ*). X-gal staining on kidney sections demonstrated patchy expression of OXGR1 in cortical tubules (Supplemental Figure 1, A and B; supplemental material available online with this article; doi:10.1172/JCI67562DS1). Costaining with a panel of cell type-specific markers revealed that OXGR1 colocalizes at the cellular level with pendrin, a $\text{Cl}^-/\text{HCO}_3^-$ exchanger that is exclusively expressed at the apical membrane of the type B and non-A–non-B intercalated cells of the CNT and the CCD (Figure 1). This specific cellular localization was validated by the absence of costaining with parvalbumin (a marker of the early distal convoluted tubule; Supplemental Figure 1, C–E), the sodium chloride cotransporter (NCC) (a marker of the early and late distal convoluted tubule; Supplemental Figure 1, F–H), the β subunit of the amiloride-sensitive epithelial sodium channel (ENaC) (a marker of the late distal convoluted tubule and the principal cells of the CNT and the CCD; Supplemental Figure 1, I–N), aquaporin-2 (a marker of principal cells of the CNT and the CCD; Supplemental Figure 1, O–T), and anion exchanger 1 (AE1) (a marker of type A intercalated cells of the CNT and the CCD; Supplemental Figure 1, U–Z). The type B and non-A–non-B intercalated cells in the CNT and the CCD are involved in the control of systemic acid-base homeostasis, extracellular fluid volume, and blood pressure (reviewed in refs. 8–10). Pendrin participates in these functions by controlling HCO_3^- secretion and electroneutral $\text{Na}^+\text{-Cl}^-$ reabsorption (upon functional coupling with Na^+ -driven $\text{Cl}^-/\text{HCO}_3^-$ exchanger NDCBE (Slc4a8) (11) and by regulating Na^+ reabsorption in the adjacent principal cells (12).

To examine whether α KG is capable of regulating apical $\text{Cl}^-/\text{HCO}_3^-$ exchange activity, we determined the activity of pendrin in intercalated cells of isolated CCDs by measuring the rate of intracellular acidification induced by the addition of luminal Cl^- (forward activity) or alkalization following Cl^- removal (reverse activity). As shown in Figure 2A, addition of 1 mM α KG to the luminal perfusion solution increased the rate of $\text{Cl}^-/\text{HCO}_3^-$ exchange in intercalated cells of CCDs from *Oxgr1*^{+/+} mice but not *Oxgr1*^{-/-}

mice, although the exchanger displayed a similar baseline activity in the two mouse strains. A representative trace depicting the effect of 1 mM α KG on intracellular pH in intercalated cells of the CCD is shown in Supplemental Figure 2. Western blot analysis demonstrated that pendrin protein abundance was not different between *Oxgr1*^{+/+} and *Oxgr1*^{-/-} mice (Supplemental Figure 3, A and D).

Next, we evaluated whether stimulation of luminal $\text{Cl}^-/\text{HCO}_3^-$ exchange was associated with increased electroneutral NaCl reabsorption. Using in vitro microperfusion of CCDs from *Oxgr1*^{+/+} mice, we found that addition of 1 mM α KG to the tubular lumen induced an equimolar reabsorption of Na^+ and Cl^- whereas basolateral α KG had no effect (Figure 2B). α KG-induced reabsorption of NaCl was markedly reduced by hydrochlorothiazide, an inhibitor of the electroneutral component of sodium reabsorption in the CCD. Interestingly, CCDs from *Oxgr1*^{-/-} mice displayed baseline Na^+ but not Cl^- transport, which was abolished by the ENaC inhibitor amiloride. Luminal addition of α KG did not increase electroneutral NaCl reabsorption in CCDs from *Oxgr1*^{-/-} mice but abolished ENaC-mediated Na^+ reabsorption (Figure 2B). Western blot analysis revealed an increase in abundance of α ENaC (*Oxgr1*^{+/+}: 100 ± 7 arbitrary units [SEM, $n = 6$] vs. *Oxgr1*^{-/-}: 136 ± 5 arbitrary units [SEM, $n = 6$], $P = 0.003$; Supplemental Figure 3, B and D) but not in abundance of β ENaC subunit (Supplemental Figure 3, C and D).

Altogether, these results demonstrate that activation of luminal OXGR1 by α KG stimulates pendrin activity and electroneutral reabsorption of NaCl by type B and non-A–non-B intercalated cells. They also suggest that ENaC activity is increased in the knockout mice and that luminal α KG may inhibit ENaC independently of OXGR1.

Next, we assessed whether challenges of acid-base or NaCl homeostasis cause changes in the luminal levels of α KG in the CNT/CCD or in the systemic circulation in mice. Wild-type C57BL/6 mice housed in metabolic cages were challenged with an oral acid, alkali, or NaCl load (0.3 M NH_4Cl , 0.28 M NaHCO_3 , or 0.28 M NaCl in drinking water, respectively) for 1, 3, 5, or 7 days. α KG levels were measured in 24-hour urine samples or in serum samples collected at the end of the experiment. As shown in Figure 3A and Supplemental Figure 4A, the acid load induced a striking decrease in urinary [α KG]/[creatinine] ratios and urinary α KG con-

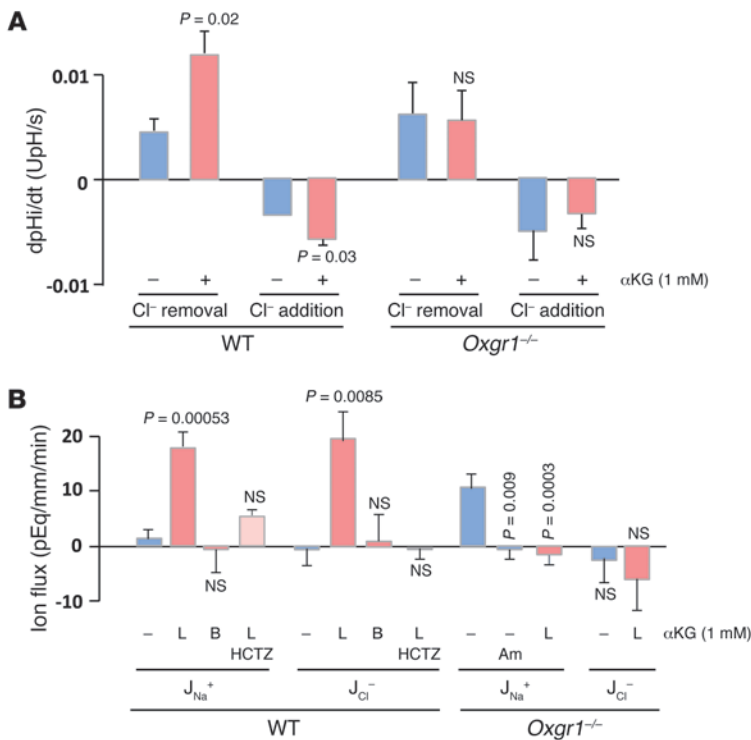


Figure 2

Activation of OXGR1 stimulates pendrin and induces NaCl reabsorption in CCD. **(A)** Apical Cl⁻/HCO₃⁻ exchange activity was measured in intercalated cells from CCDs isolated from wild-type and *Oxgr1*^{-/-} mice in the absence or the presence of 1 mM αKG in the luminal fluid. The exchanger activity was determined either in the forward (Cl⁻ addition induces intracellular acidification) or reverse direction (Cl⁻ removal induces intracellular alkalization). Values are mean ± SEM from 3 CCDs, and the exchange activity was measured on at least 3 cells per CCD. dpHi/dt, initial rate of change in cell pH. *P* values are shown in comparison with controls (absence of αKG) (Student's *t* test). **(B)** Na⁺ and Cl⁻ reabsorption fluxes (*J*_{Na⁺} and *J*_{Cl⁻}, respectively) measured in vitro—microperfused CCDs from wild-type and *Oxgr1*^{-/-} mice were determined in the absence or presence of 1 mM αKG in the luminal (L) or peritubular (B) solution. Some CCDs were pretreated with 100 μM hydrochlorothiazide (HCTZ) or 10 μM amiloride (Am). Note that in absence of αKG, CCDs from wild-type mice displayed no NaCl transport whereas those from *Oxgr1*^{-/-} mice showed amiloride-sensitive Na⁺ reabsorption that was inhibited upon addition of αKG. Values are mean ± SEM from 4 to 6 CCDs. *P* values are shown in comparison with controls (absence of αKG) (variance analysis [ANOVA]).

centrations, respectively, whereas base load resulted in an opposite pattern, with a significant increase in urinary αKG levels. The changes in urinary αKG levels were parallel to changes in urinary pH (Supplemental Figure 4B). At the same time, addition of NaCl to drinking water did not produce any effect on urinary αKG levels (Figure 3A and Supplemental Figure 4A). Importantly, none of the dietary supplements had a significant effect on plasma αKG concentration, even though there was a tendency for increased αKG concentration in mice loaded with NaHCO₃ (Figure 3B). To test whether base load can produce more rapid (<24 hours) effects on urinary αKG levels and to assess the threshold dose of base capable of increasing αKG concentrations in urine, mice were gavaged with 400 μl solution containing various concentrations of NaHCO₃. Spot urine was collected 3 hours after gavage. As control groups, mice gavaged with 400 μl of either water or 0.56 M NaCl were used. As shown in Figure 3C, mice gavaged with water or 0.56 M NaCl excreted similar amounts of αKG in the urine. Base load induced a significant increase in both urinary pH and αKG/creatinine ratios when compared with those in mice gavaged with water or NaCl controls (Figure 3C). Importantly, these changes reached statistical significance already at low doses of NaHCO₃ (22.4 μmol NaHCO₃ for αKG/creatinine ratio and 44.8 μmol NaHCO₃ for urinary pH). Collectively, these results demonstrated that the challenge of mice with acid-base but not NaCl load can produce rapid and significant effects on urinary αKG levels.

To test whether deficiency of OXGR1 could affect acid-base equilibrium, we analyzed urine and blood samples of wild-type and *Oxgr1*^{-/-} mice. As shown in Table 1, *Oxgr1*^{-/-} mice fed standard diet had a small but statistically significant decrease in urine pH and a statistically significant increase in urinary titratable acid and αKG levels. These results suggested that the acid-base equilibrium might be disturbed in *Oxgr1*^{-/-} mice. However, blood chemistry parameters were not different between wild-type and *Oxgr1*^{-/-} mice (Table 1).

Thus, we tested the role of OXGR1 in the condition of experimentally induced alkalosis. As shown in Table 1, both wild-type and *Oxgr1*^{-/-} mice given a NaHCO₃-rich diet for 3 days (see Methods) developed a metabolic alkalosis with plasma HCO₃⁻ levels raised above 30 mM. However, the increase in plasma HCO₃⁻ levels was significantly higher in *Oxgr1*^{-/-} mice. Tubular handling of bicarbonate under alkali load was examined in a separate experiment by assessing its fractional excretion (FE_{HCO₃}). As shown in Table 1, the FE_{HCO₃} was more than 2-fold lower in *Oxgr1*^{-/-} mice compared with that in wild-type mice, thus suggesting an inappropriate renal/tubular response in *Oxgr1*^{-/-} mice to the alkali load. Collectively, these results indicated that renal mechanisms responsible for maintaining acid-base equilibrium upon alkali load are impaired in *Oxgr1*^{-/-} mice. The fact that plasma Cl⁻ concentration was significantly lower in *Oxgr1*^{-/-} mice (Table 1) suggested that a decrease in the HCO₃⁻/Cl⁻ exchange activity via pendrin is involved in this impairment.

Discussion

Paracrine signaling plays an important role in renal function. For instance, the tubuloglomerular feedback response, which is mediated by a multitude of paracrine molecules (ATP, adenosine, prostaglandin E2, and nitric oxide), is critically involved in the regulation of glomerular filtration rate (reviewed in ref. 13). In this study, we show for the first time that maintaining systemic acid-base balance requires paracrine communication between the proximal and the distal parts of the renal tubule. We demonstrate that αKG acts in a paracrine manner on type B and non-A-non-B intercalated cells of the CNT/CCD, stimulating HCO₃⁻ secretion and NaCl reabsorption. These effects are induced by apical but not basolateral application of αKG and are mediated by OXGR1, a GPCR that is specifically expressed in type B and non-A-non-B intercalated cells. Importantly, our estimations suggest that luminal levels of αKG in the CNT/CCD vary within the range of αKG

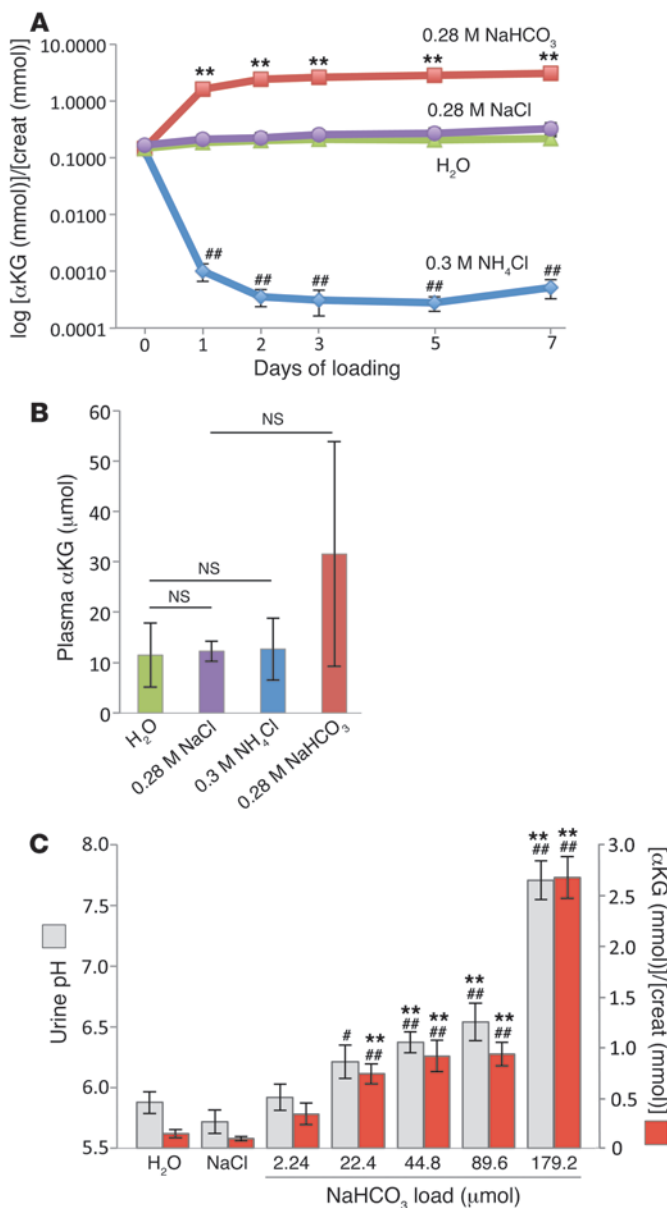


Figure 3

Effects of a dietary load of acid, base, or NaCl on urinary and plasma levels of α KG. **(A)** Analysis of $[\alpha\text{KG}]/[\text{creatinine}]$ ratios in urine collected from wild-type mice housed individually in metabolic cages. The 24-hour urine was collected from 4 groups of mice (6 mice per group): a control group receiving tap water (green) and groups supplemented with acid (0.3 M NH_4Cl ; blue), base (0.28 M NaHCO_3 ; red), or salt (0.28 M NaCl ; purple) in drinking water. Values are mean \pm SEM. ** $P < 0.005$, compared with 0.28 M NaCl group; ### $P < 0.005$, compared with H_2O group; Student's t test. **(B)** Analysis of plasma α KG concentrations in wild-type mice loaded for 7 days with acid (0.3 M NH_4Cl ; blue), base (0.28 M NaHCO_3 ; red), or salt (0.28 M NaCl ; purple) in drinking water. Mice in the control group received tap water (green). Values are mean \pm SD ($n = 6-9$); Student's t test. **(C)** Analysis of urinary $[\alpha\text{KG}]/[\text{creatinine}]$ ratios (red bars) and urinary pH (gray bars) in spot urine of wild-type mice gavaged with 400 μl of tap water, 400 μl of 0.56 M NaCl , or 400 μl of NaHCO_3 at different concentrations (0.0056 M, 0.056 M, 0.112 M, 0.224 M, and 0.448 M). The urine was collected 3 hours after gavage. Values are mean \pm SEM. ** $P < 0.005$, compared with water group; ### $P < 0.005$, compared with 0.56 M NaCl group; # $P < 0.01$, compared with controls (variance analysis [ANOVA]).

to acid-base stress and that these changes can be induced by low, physiologically relevant doses of acid or base (e.g., the dose of 22.4 μmol NaHCO_3 , which produced a significant effect on the urinary α KG levels in gavage experiments, is significantly lower than the amount of HCO_3^- produced by the metabolism of 1 g of the 24% protein-containing standard diet used in this study). This suggests that the activation level of this paracrine mechanism can be modulated by the diurnal variations in the dietary acid-base load and/or by the variations in body's metabolic production of acid and base.

We propose that α KG/OXGR1 paracrine signaling is involved in maintenance of acid-base balance at least in two ways. First, this paracrine mechanism controls pendrin activity, thereby exerting a direct effect on the acid-base balance. The rapid onset of pendrin activation by α KG/OXGR1 (within 20 minutes) suggests that this regulation is transcriptionally independent. Together with the results showing rapid changes in urinary α KG levels under acute acid-base stress, these observations support the hypothesis that the α KG/OXGR1/pendrin axis is involved in the dynamic fine-tuning of bicarbonate secretion by type B and non-A-non-B intercalated cells. Second, we hypothesize that α KG/OXGR1 signaling may be involved in the compensatory response to the acid-base stress-induced changes in NaCl reabsorption through the NHE3 in the proximal tubule and Henle's loop. Indeed, it was demonstrated that acute acid or base loading induces an increase or decrease in the NHE3 activity, respectively (reviewed in ref. 14). Because NHE3 is functionally coupled to a Cl^- anion exchange, these changes result not only in increased or decreased bicarbonate reabsorption but also in increased or decreased net NaCl reabsorption in the proximal tubule, respectively. Thus, by decreasing or increasing NaCl reabsorption in the type B and non-A-non-B intercalated cells under acid or base loading, respectively, the α KG/OXGR1 signaling pathway participates in maintaining NaCl homeostasis and, indirectly, in acid-base balance. This model is depicted in Supplemental Figure 5. To what extent this paracrine mechanism operates similarly/differentially between type B and type non-A/non-B intercalated

concentrations that have been shown to activate OXGR1 in vitro ($\text{EC}_{50} \sim 70 \mu\text{M}$; ref. 1). Indeed, considering that under our experimental condition urine was concentrated approximately 10–20 times between the lumen of the CNT/CCD and the final urine, one can estimate the range of α KG concentration at the site of OXGR1 expression as several hundreds nanomolar under acid load, as one hundred to several hundreds micromolar under standard diet, and as one to several millimolar under base load (e.g., the concentration of α KG in the final urine of mice given 0.3 M NH_4Cl in drinking water for 2 days is $3.53 \pm 2.10 \mu\text{M}$ [$n = 6$, SD], the concentration of α KG in the final urine of mice given tap water is $2.03 \pm 0.67 \text{ mM}$ [$n = 6$, SD], and the concentration of α KG in the final urine of mice given 0.28 M NaHCO_3 in drinking water for 2 days is $36.43 \pm 8.16 \text{ mM}$ [$n = 6$, SD]; see Supplemental Figure 4A). Another piece of evidence supporting the functional relevance of this paracrine signaling is the finding that significant changes in urinary levels of α KG occur rapidly (less than 3 hours) in response

**Table 1**
Blood and urine chemistry in wild-type and *Oxgr1*^{-/-} mice

Standard diet	Wild-type	<i>Oxgr1</i> ^{-/-}	P
Blood/plasma			
Na ⁺ (mM)	151.8 ± 3.1 (7)	146.1 ± 1.6 (7)	NS
K ⁺ (mM)	4.6 ± 0.2 (7)	4.6 ± 0.1 (7)	NS
Cl ⁻ (mM)	118.7 ± 2.0 (10)	116.9 ± 2.4 (10)	NS
HCO ₃ ⁻ (mM)	22.6 ± 0.4 (10)	22.8 ± 0.5 (10)	NS
pCO ₂ (mmHg)	45.0 ± 4.2 (10)	38.6 ± 1.4 (10)	NS
pO ₂ (mmHg)	128.0 ± 10 (10)	127.5 ± 7.0 (10)	NS
pH	7.35 ± 0.02 (10)	7.39 ± 0.02 (10)	NS
Osmolality (mOsm)	325.6 ± 4.6 (5)	324.3 ± 1.2 (7)	NS
Aldosterone (pg/ml)	337.5 ± 36.9 (11)	380.5 ± 26.1.2 (10)	NS
Urine			
Na ⁺ /creatinine (mmol/mmol)	11.9 ± 0.4 (32)	12.0 ± 0.5 (33)	NS
K ⁺ /creatinine (mmol/mmol)	52.7 ± 1.5 (32)	57.4 ± 2.5 (33)	NS
Cl ⁻ /creatinine (mmol/mmol)	20.1 ± 1.1 (23)	22.7 ± 1.4 (25)	NS
TA/creatinine (mmol/mmol)	6.1 ± 0.4 (23)	8.0 ± 0.6 (25)	0.008
NH ₄ ⁺ /creatinine (mmol/mmol)	11.8 ± 1.1 (23)	12.6 ± 1.4 (25)	NS
αKG/creatinine (mmol/mmol)	0.11 ± 0.01 (23)	0.14 ± 0.01 (25)	0.03
pH	6.24 ± 0.02 (32)	6.10 ± 0.02 (33)	0.00001
Volume (ml/24 h)	1.36 ± 0.07 (32)	1.41 ± 0.07 (33)	NS
Base-rich diet (3 days)			
Blood/plasma			
Na ⁺ (mM)	147.9 ± 0.6 (10)	147.9 ± 1.0 (10)	NS
Cl ⁻ (mM)	113.7 ± 0.7 (10)	112.0 ± 1.0 (10)	0.0004
HCO ₃ ⁻ (mM)	30.6 ± 1.5 (10)	33.4 ± 2.89 (10)	0.01
pH	7.38 ± 0.06 (10)	7.39 ± 0.04 (10)	NS
Urine			
Na ⁺ /creatinine (mmol/mmol)	57.1 ± 3.6 (6)	53.2 ± 9.3 (6)	NS
K ⁺ /creatinine (mmol/mmol)	14.0 ± 1.1 (6)	13.3 ± 3.0 (6)	NS
Cl ⁻ /creatinine (mmol/mmol)	9.9 ± 0.5 (6)	9.2 ± 2.1 (6)	NS
pH	9.00 ± 0.11 (6)	8.95 ± 0.17 (6)	NS
Volume (ml/24 h)	4.84 ± 0.40 (6)	5.23 ± 1.47 (6)	NS
FE _{HCO3} (%)	3.5 ± 1.4 (5)	1.5 ± 0.5 (5)	0.017

Values are mean ± SD (Student's *t* test). Numbers in parentheses represent the number of mice used to determine each parameter. All urine and blood samples were collected from individual mice.

cells or between the CNT and CCD (which contain different ratios of these cell types) remains to be established in future studies. The idea that the αKG/OXGR1 pathway is involved in sodium balance is supported by the fact that *Oxgr1*^{-/-} mice exhibit an increased functional activity of ENaC, correlating with the increased protein abundance of αENaC subunit. The mechanism of these adaptive changes remains unknown, but our data indicate that they occur without changes in plasma aldosterone levels.

Besides its OXGR1-mediated stimulatory action on NaCl reabsorption in intercalated cells, αKG also inhibits amiloride-sensitive Na⁺ reabsorption in principal cells independently of OXGR1 activation. This metabolic effect of αKG may be related to increased ATP production and inhibition of ENaC via autocrine activation of P2Y2 receptors (15). Receptor-dependent and -independent effects of αKG on intercalated and principal cells both converge to compensate for alkalosis-induced decrease in proximal tubule reabsorption of NaCl by favoring NaCl reabsorption over Na⁺/K⁺ exchange along the CNT/CCD.

Our finding that αKG serves as a paracrine mediator in regulating acid-base balance adds to the accumulating evidence for the

regulatory homeostatic role of organic anions excreted in urine. It has been long recognized that citrate, the major intermediate of TCA cycle in the urine, has a dual role in maintaining acid-base balance and in the chelation of calcium ions, thereby preventing calcium stone formation (reviewed in ref. 16). Peti-Peterdi and colleagues have demonstrated recently the role of luminal succinate in GPR91-mediated regulation of renin secretion (17, 18). Today, citrate supplementation is widely used in the prevention of nephrolithiasis and GPR91 is recognized as a potential target in treatment of diabetic nephropathy. Our results and the fact that OXGR1 expression is almost exclusively restricted to the kidney suggest that OXGR1 could represent an interesting therapeutic opportunity for treatment of acid-base disorders or in the treatment of tubular disorders associated with kidney stone formation. The second tissue that exhibits a high level of OXGR1 expression is the testis and epididymis (19). Whether OXGR1 is involved in the homeostasis of epididymal fluid and male fertility needs to be addressed by future investigation.

Methods

Animals. A colony of *Oxgr1*^{-/-} mice (C57BL/6J background) was established from breeding pairs of *Oxgr1*^{+/-} heterozygous mice developed by VelociGene and obtained from the Knock-Out Mouse Project repository. Male mice weighing 25–30 g (8 weeks of age) were used in all experiments. The animals were maintained on the standard laboratory chow diet (KLIBA NAFAG diet 3800). The NaHCO₃-rich and standard jelly diets were prepared as follows: 6 ml 0.7 M NaHCO₃ or 6 ml water were mixed with 0.2 ml agar-agar solution and briefly boiled. The resulting solution was cooled down to 40°C and mixed with 5 g of the standard lab chow in the form of powder. The final mixture was stored at 4°C for a maximum of 3 days. The mice received 11 g of the jelly food per day. All experiments with animals were performed in accordance with the Swiss guidelines for animal care.

Antibodies. The antibodies against NCC, aquaporin-2, αENaC, βENaC, AE1, and pendrin have been used as previously described (20, 21). Anti-parvalbumin antibody was from Swant. Anti-β-galactosidase antibody was from Abcam. Anti-actin antibody was from Sigma-Aldrich. We have tested several commercially available anti-OXGR1 antibodies (2-oxoglutarate receptor 1 antibody 18-461-10700 from GenWay, anti-OXGR1 antibody ab67351 from Abcam, and anti-OXGR1 antibody LS-A1865 from LifeSpan BioSciences); however, none of them have shown specific results. Immunohistochemistry and X-gal staining protocols are available in the Supplemental Methods.

Metabolic cages. Mice were housed in individual metabolic cages (Tecniplast). Urine collection was performed after a 3-day adaptation period. Urine osmolality, pH, and ionic composition were analyzed as previously described (22).

Blood sample analysis. In experiments with alkali load, blood pH and blood gases were analyzed in blood samples collected from the retro-orbital sinus under isoflurane anesthesia. Blood was collected through a catheter implanted into the left carotid artery 4 hours before blood collection from mice fed a normal diet.



α KG measurement. Urine and plasma α KG were measured using an Alpha-Ketoglutarate Assay Kit from BioVision.

Ion flux in isolated CCD. CCDs from wild-type and *Oxgr1*^{-/-} mice were microdissected from corticomedullary rays and microperfused under symmetrical conditions. The bath and perfusate contained 118 mM NaCl, 23 mM NaHCO₃, 1.2 mM MgSO₄, 2 mM K₂HPO₄, 2 mM calcium lactate, 1 mM Na-citrate, 5.5 mM glucose, and 12 mM creatinine, pH 7.4 (bath continuously gassed with 95% O₂/5% CO₂) at 37°C. Measurements were conducted during the first 90 minutes of perfusion. Hydrochlorothiazide and amiloride were added to the perfusate at a final concentration of 10⁻⁴ M and 10⁻⁵ M, respectively. Collections from 4 periods of 15 minutes were performed in which 30–35 nl of fluid was collected. The collection volume was determined under water-saturated mineral oil with calibrated volumetric pipettes. Concentrations of Na⁺ and creatinine were determined by HPLC using 26 nl of collected fluid. Concentrations of Cl⁻ were measured by microcoulometry on 2 to 3 nl of collected fluid. For each collection, the ion flux of sodium (J_x) was calculated per unit length of tubule: $J_x = \{([X]_p \times V_p) - ([X]_c \times V_c)\} / (L \times t)$, where $[X]_p$ and $[X]_c$ are the concentrations of X in the perfusate and collection, respectively; V_p and V_c are the perfusion and collection rates, respectively; L is the tubule length; and t is the collection time. V_p was calculated as $V_p = V_c \times [\text{creat}]_c / [\text{creat}]_p$, where $[\text{creat}]_c$ and $[\text{creat}]_p$ are the concentrations of creatinine in the collection and perfusate, respectively. For each tubule, ion fluxes were calculated as the mean of the 4 collection periods.

Apical Cl⁻/HCO₃⁻ exchange activity in CCD intercalated cells. The activity of pendrin was determined by measuring the rate of intracellular alkalinization in type B and non-A-non-B intercalated cells following removal of luminal chloride as previously described (see Supplemental Methods for cell type identification). Our setup allowed a smooth and complete exchange of the luminal fluid within 3 to 4 seconds. Briefly, CCDs were dissected and microperfused in vitro as described above. Intracellular pH was assessed with imaging-based, dual excitation wavelength fluorescence microscopy using 2',7'-Bis-(2-Carboxyethyl)-5-(and-6)-carboxyfluorescein (BCECF, Molecular Probes). Intracellular dye was excited alternatively at 440 and 500 nm every 2 seconds with a light-emitting diode (Optoled, Cairn Research). Emitted light was collected through a dichroic mirror, passed through a 530-nm filter, and focused onto a EM-CCD camera (iXon, Andor Technology) connected to a computer. The measured light intensities were digitized with 14-bit precision (16,384-gray-level scale) for further analysis. For each tubule, 3–4 intercalated cells were analyzed and the mean gray level was measured

with Andor IQ software (Andor Technology). Background fluorescence was subtracted from fluorescence intensity to obtain intensity of intracellular fluorescence. Intracellular dye was calibrated at the end of each experiment using the high [K⁺]-nigericin technique. CCDs were initially perfused with a Cl⁻-containing solution (119 mM NMDG-Cl, 23 mM NMDG-HCO₃, 2 mM K₂HPO₄, 1.5 mM CaCl₂, 1.2 mM MgSO₄, 10 mM HEPES, and 5.5 mM D-glucose) and then switched to a Cl⁻-free solution (119 mM NMDG-glucuronate, 23 mM NMDG-HCO₃, 2 mM K₂HPO₄, 7.5 mM Ca-gluconate, 1.2 mM MgSO₄, 10 mM HEPES, and 5.5 mM D-glucose). All solutions were adjusted to pH 7.40 and continuously bubbled with 95% O₂/5% CO₂.

Statistics. Data are presented as mean \pm SEM or as mean \pm SD, as indicated in the corresponding figure legends. The tests used include the 2-tailed, unpaired t test and the 1-way ANOVA. A P value of less than 0.05 was considered significant.

Study approval. Animal studies were approved by the Veterinary Service of the Canton de Vaud, Switzerland.

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